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(FILE 'HOME' ENTERED AT 09:52:14 ON 03 JUN 2003)

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FILE 'CAPLUS, BIOSIS, SCISEARCH, LIFESCI, PASCAL, EMBASE, MEDLINE, ESBIOBASE, AQUASCI, BIOTECHNO' ENTERED AT 09:54:09 ON 03 JUN 2003 27 S L1 AND (PROMOTER OR REPORTER ENZYME OR REPORTER SYSTEM) 8 DUP REM L2 (19 DUPLICATES REMOVED)

=> log Y

L1

L2L3 => d 13 ibib ab 1-8

L3 ANSWER 1 OF 8 CAPLUS COPYRIGHT 2003 ACS ACCESSION NUMBER: 2002:107593 CAPLUS

DOCUMENT NUMBER:

136:161320

TITLE:

Use of ectoenzymes and secreted enzymes to monitor

cellular proliferation

INVENTOR (S):

Zyskind, Judith

PATENT ASSIGNEE(S):

Elitra Pharmaceuticals, Inc., USA

SOURCE:

PCT Int. Appl., 80 pp. CODEN: PIXXD2

DOCUMENT TYPE:

Patent

LANGUAGE:

English

FAMILY ACC. NUM. COUNT:

r: 1

PATENT INFORMATION:

PATENT NO. KIND DATE APPLICATION NO. DATE ---------------WO 2002010442 A1 20020207 WO 2000-US21049 20000802 W: AE, AG, AL, AM, AT, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CR, CU, CZ, CZ, DE, DE, DK, DK, DM, DZ, EE, EE, ES, FI, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ,

CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
PRIORITY APPLN. INFO.: WO 2000-US21049 20000802

The invention concerns a method for measuring cellular proliferation in a sample comprising obtaining a sample of cells which express an ectoenzyme or a secreted enzyme, detg. the level of activity of the ectoenzyme or secreted enzyme in the sample and correlating the level of activity of the ectoenzyme or secreted enzyme with the extent of cellular proliferation. Further, secreted enzymes and ectoenzymes such as membrane-bound chitobiase (N,N'-diacetylchitobiase) and nucleic acids are used in combination with genetic methods to det. the impact of test compds. on cell proliferation.

REFERENCE COUNT:

5 THERE ARE 5 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L3 ANSWER 2 OF 8 CAPLUS COPYRIGHT 2003 ACS

ACCESSION NUMBER:

2001:284144 CAPLUS

DOCUMENT NUMBER:

INVENTOR(S):

134:306156

TITLE:

A cytoplasmic form of chitobiase as a

reporter enzyme for the study of

gene expression Zyskind, Judith

PATENT ASSIGNEE(S):

Elitra Pharmaceuticals, Inc., USA

SOURCE:

PCT Int. Appl., 44 pp.

CODEN: PIXXD2

DOCUMENT TYPE:

Patent English

LANGUAGE:

r. 1

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 2001027322	A2	20010419	WO 2000-US21048	20000802
WO 2001027322	A3	20011213		

W: AE, AG, AL, AM, AT, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CR, CU, CZ, CZ, DE, DE, DK, DK, DM, DZ, EE, EE, ES, FI, FI,

GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG N. INFO.: US 1999-159221P P 19991013 PRIORITY APPLN. INFO.: The present invention relates to reporter gene constructs encoding a cytoplasmic form of chitobiase (N,N'-diacetylchitobiase) and methods of using these reporter gene constructs. The use of a cytoplasmic form of chitobiase as a reporter enzyme is generally applicable to the study of gene expression in organisms which do not contain N-acetyl-.beta.-D-glucosaminidases. ANSWER 3 OF 8 CAPLUS COPYRIGHT 2003 ACS ACCESSION NUMBER: 2000:34968 CAPLUS DOCUMENT NUMBER: 132:89234 Fungal signal peptide and their use in secreting chitinolytic proteins from transgenic plants INVENTOR(S): Harman, Gary E.; Lorito, Matteo; Woo, Sheridan; Brants, Aigars; Earle, Elizabeth; Kubicek, Christian P.; Peterbauer, Clemens K.; Tronsmo, Arne; Klemsdahl, Sonja PATENT ASSIGNEE(S): Cornell Research Foundation, Inc., USA PCT Int. Appl., 88 pp. SOURCE: CODEN: PIXXD2 DOCUMENT TYPE: Patent. LANGUAGE: English FAMILY ACC. NUM. COUNT: PATENT INFORMATION: PATENT NO. KIND DATE APPLICATION NO. DATE -----

TITLE:

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    WO 2000001812
                     A1
                           20000113
                                         WO 1999-US15242 19990706
        W: AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE,
            DK, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP,
            KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN,
            MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM,
            TR, TT, UA, UG, UZ, VN, YU, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ,
        RW: GH, GM, KE, LS, MW, SD, SL, SZ, UG, ZW, AT, BE, CH, CY, DE, DK,
            ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG,
            CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
    AU 9948621
                                         AU 1999-48621
                     A1
                         20000124
                                                          19990706
PRIORITY APPLN. INFO.:
                                       US 1998-91768P P 19980706
                                       WO 1999-US15242 W 19990706
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AB Signal peptides from secreted proteins of fungi can be used to direct secretion of proteins from plant cells. Transgenic plants contg. expression constructs using these signal peptides can be regenerated. Specifically, signal peptides from chitinases of Trichoderm harzianum P1 are described. The chitinases can be used to improve plant resistance to insect pests and phytopathogenic fungi. An endochitinase and a chitobiase from cultures of T. harzianum were shown to inhibit fungal spore germination and to inhibit growth of insect larvae. A cDNA for the endochitinase was cloned by screening an expression library with antibodies to the enzyme. Similarly, a chitinase from Gliocladium virens was also found to have fungicidal effects. The T. harzianum chitinase cDNA was placed under control of a 35S promoter and introduced into tobacco and potato. Transgenic plants transcribed the gene and synthesized the protein with endogenous chitinase levels increased 10-400-fold. Transgenic plants showed greatly increased resistance to challenge with the airborne pathogen Alternaria alternata with some lines being completely resistant. Similarly, they also showed increased

resistance to the soil-borne Rhizoctonia solani.

REFERENCE COUNT:

THERE ARE 3 CITED REFERENCES AVAILABLE FOR THIS 3

RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

ANSWER 4 OF 8 SCISEARCH COPYRIGHT 2003 THOMSON ISIDUPLICATE 1

1999:288244 SCISEARCH ACCESSION NUMBER:

THE GENUINE ARTICLE: 184QP

Multiple genes involved in chitin degradation from the TITLE:

marine bacterium Pseudoalteromonas sp. strain S91

AUTHOR: Techkarnjanaruk S; Goodman A E (Reprint)

UNIV S AUSTRALIA, SCH BIOL SCI, GPO BOX 2100, ADELAIDE, SA CORPORATE SOURCE:

5001, AUSTRALIA (Reprint); UNIV S AUSTRALIA, SCH BIOL SCI,

ADELAIDE, SA 5001, AUSTRALIA

COUNTRY OF AUTHOR: AUSTRALIA

SOURCE: MICROBIOLOGY-UK, (APR 1999) Vol. 145, Part 4, pp. 925-934.

Publisher: SOC GENERAL MICROBIOLOGY, MARLBOROUGH HOUSE, BASINGSTOKE RD, SPENCERS WOODS, READING RG7 1AE, BERKS,

ENGLAND.

ISSN: 1350-0872. Article; Journal

DOCUMENT TYPE: FILE SEGMENT:

LIFE

LANGUAGE:

English

REFERENCE COUNT:

ABSTRACT IS AVAILABLE IN THE ALL AND IALL FORMATS

A cluster of three closely linked chitinase genes organized in the order chiA, chiB and chiC, with the same transcriptional direction, and two unlinked genes, chiP and chiB involved in chitin degradation in Pseudoalteromnas sp. strain S91 were cloned, sequenced and characterized. The deduced amino acid sequences revealed that ChiA, ChiB and ChiC exhibited similarities to chitinases belonging to family 18 of the glycosyl hydrolases while ChiP and ChiQ belonged to family 20. ChiP and ChiQ showed different enzymic activities against fluorescent chitin analogues, but neither was able to degrade colloidal chitin. ChiA possessed chitinase activity but did not bind chitin; ChiB bound chitin but had no chitinase activity; ChiC possessed strong chitinase activity and also bound chitin. Production of ChiC in S91 appeared to be controlled by chiA expression, since insertion of a transposon into the ORF of chiA resulted in the loss of chitinase activity as well as loss of ChiC proteins in a chitinase-negative mutant. In Escherichia coli, ChiC appeared to be expressed from its own promoter.

ANSWER 5 OF 8 CAPLUS COPYRIGHT 2003 ACS DUPLICATE 2

ACCESSION NUMBER: 1999:336322 CAPLUS

DOCUMENT NUMBER: 131:154270

TITLE: Structure of the human gene for lysosomal

di-N-acetylchitobiase

AUTHOR(S):

CORPORATE SOURCE:

Liu, Bei; Ahmad, Wasim; Aronson, Nathan N., Jr. Department of Biochemistry and Molecular Biology,

College of Medicine, University of South Alabama,

Mobile, AL, 36688, USA

SOURCE: Glycobiology (1999), 9(6), 589-593

CODEN: GLYCE3; ISSN: 0959-6658

PUBLISHER:

Oxford University Press

DOCUMENT TYPE:

Journal

LANGUAGE:

English

Chitobiase is a lysosomal glycosidase that acts during the ordered degrdn. of asparagine-linked glycoproteins to cleave the core chitobiose unit at its reducing end. Human chitobiase is expressed in significant amts., while bovine chitobiase is produced at extremely low levels. To begin to understand this

species-dependent expression, the authors detd. the gene structure of

human chitobiase. The human chitobiase gene (CTBS) is

approx. 20 kb comprising seven exons varying from 0.1 to 2.3 kb and six introns of 0.3 to 8 kb. The previously characterized partial bovine

chitobiase gene structure is similarly organized including exon and intron sizes and locations, but the human and bovine 5'-flanking regions differ significantly. 5'-RACE anal. of human chitobiase cDNA revealed only one transcriptional start site 45 bp upstream of the ATG translation initiation site. Computer anal. of the human chitobiase gene 5'-flanking region shows characteristics of a typical housekeeping gene. The putative promoter region contains a distal TATA box, and there are several Sp-1 and AP-2 cis elements. In contrast, bovine chitobiase gene 5'-flanking region shows totally different structures and may contain several silencers. A partial art-2 segment which is an artiodactyl Alu-like repetitive sequence, is also present. These evolutionary differences in the 5'-flanking region of the chitobiase genes from human and bovine could account for the widely varied expression levels of the hydrolase within these two species.

REFERENCE COUNT: 18

THERE ARE 18 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

ANSWER 6 OF 8 CAPLUS COPYRIGHT 2003 ACS

DUPLICATE 3

ACCESSION NUMBER:

1999:3059 CAPLUS 130:149244

DOCUMENT NUMBER: TITLE:

Chitobiase, a new reporter

enzyme AUTHOR (S):

Kalabat, D. Y.; Froelich, J. M.; Phuong, T. K.;

Forsyth, R. A.; Newman, V. G.; Zyskind, J. W. San Diego State University, San Diego, CA, USA

CORPORATE SOURCE: SOURCE:

BioTechniques (1998), 25(6), 1030-1035 CODEN: BTNQDO; ISSN: 0736-6205

PUBLISHER:

Eaton Publishing Co.

DOCUMENT TYPE: LANGUAGE:

Journal English

N,N'-diacetylchitobiase from the marine organism Vibrio harveyi is a highly stable reporter enzyme for gene fusions. This enzyme hydrolyzes the disaccharide chitobiose to N-acetylglucosamine. advantages of the reporter gene encoding chitobiase (chb) are: (i) that chitobiase and N-acetyl-.beta.-D-glucosaminidase activities are missing in E. coli strains, (ii) chitobiase can be monitored using blue/white colony indicator plates and (iii) convenient substrates for this enzyme are com. available. The use of chitobiase as a reporter enzyme is generally applicable to the study of gene expression in those bacteria that do not contain N-acetyl-.beta.-D-glucosaminidases. The authors constructed plasmid vectors contg. a multiple cloning site for producing in-frame fusions to chitobiase, the attP of .lambda. phage for movement into the bacterial chromosome for single-copy anal., the gene encoding chloramphenicol acetyltransferase (cat), the pACYC184 origin of replication and the rrnBt1t2 terminator region upstream of the chb gene to prevent read-through from other promoters. In-frame fusions between the dnaA gene and chb were moved to the chromosome by site-specific recombination with the chromosomal attB site. These single-copy fusions were assayed for chitobiase to examine the

effects of a deletion in the dnaA regulatory region. REFERENCE COUNT: THERE ARE 19 CITED REFERENCES AVAILABLE FOR THIS 19 RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

ANSWER 7 OF 8 CAPLUS COPYRIGHT 2003 ACS

DUPLICATE 4

ACCESSION NUMBER:

1990:231809 CAPLUS

DOCUMENT NUMBER:

112:231809

TITLE:

N,N'-Diacetylchitobiase of Vibrio harveyi. Primary structure, processing, and evolutionary relationships

AUTHOR(S): CORPORATE SOURCE:

Soto-Gil, Rafael W.; Zyskind, Judith W.

Mol. Biol. Inst., San Diego State Univ., San Diego,

SOURCE:

CA, 92182, USA Journal of Biological Chemistry (1989), 264(25), 14778-83

CODEN: JBCHA3; ISSN: 0021-9258

DOCUMENT TYPE:

LANGUAGE:

Journal English

The nucleotide sequence of the gene chb, encoding the outer membrane protein, N,N'-diacetylchitobiase (chitobiase), of the marine bacterium, Vibrio harveyi, was detd. The amino acid sequence of prechitobiase was derived from the nucleotide sequence. Prechitobiase has a mol. mass of 97,771 Da and consists of 883 amino acid residues. characteristic signal peptide is at the N-terminus whose removal is inhibited by the antibiotic, globomycin, suggesting that mature chitobiase is a lipoprotein with a maturation pathway similar to that of the Escherichia coli major outer membrane lipoprotein. A perfect homol. of 6 amino acids at the processing and modification region of the outer membrane lipoprotein of E. coli was found with amino acids 15-19 of the deduced prechitobiase protein sequence. Chitobiase shares similarities and possibly common ancestry with the .alpha.-chain of the human .beta.-hexosaminidase. A comparison of the amino acid sequences of chitobiase and the .alpha.-chain of .beta.-hexosaminidase gave a highly significant alignment score of 19.1 std. deviation units above a mean randomized alignment score. Primer extension anal. of the promoter region revealed 3 transcription initiation sites used by E. coli cells harboring the chb gene, 2 of which were also evident in V. harveyi cells.

ANSWER 8 OF 8 CAPLUS COPYRIGHT 2003 ACS

DUPLICATE 5

ACCESSION NUMBER: DOCUMENT NUMBER:

1986:492435 CAPLUS 105:92435

TITLE:

Chitinase determinants of Vibrio vulnificus: gene

cloning and applications of a chitinase probe

AUTHOR(S): CORPORATE SOURCE:

Wortman, A. T.; Somerville, C. C.; Colwell, R. R. Dep. Microbiol., Univ. Maryland, College Park, MD,

20742, USA

SOURCE:

Applied and Environmental Microbiology (1986), 52(1),

142-5

CODEN: AEMIDF; ISSN: 0099-2240

DOCUMENT TYPE:

Journal

LANGUAGE:

English To initiate study of the genetic control of chitinolytic activity in AB vibrios, the chitobiase [9012-33-3] gene was isolated by

cloning chromosomal DNA prepd. from V. vulnificus. Chimeric plasmids were constructed from Sau3A I partial digests of chromosomal DNA prepd. by ligating 5-15-kilobase fragments into the BamHI site, i.e., in the tetracycline-resistance (Tcr) gene, of pBR322 (AmrTcr). The resulting plasmids were transformed into Escherichia coli DH1. Chitobiase activity of the insert-bearing clones was detected by using a chromogenic substrate, p-nitrophenyl-N-acetyl-.beta.,D-glucosaminide, and confirmed by the appearance of a fluorescent end product from the hydrolysis of 4-methylumbelliferyl-.beta.,D-N-N'-diacetylchitiobiose. Endochitinase activity was demonstrated by liberation of water-sol. products produced by the degrdn. of [3H] chitin. Transformation of E. coli Y10R (lacY) with plasmids from chitinase-pos. clones restored the lactose-pos. phenotype, suggesting the presence of a permease assocd. with chitinase activity. Phys. mapping of plasmids contg. the chitinase determinants indicate that transcription of these genes in E. coli may be initiated at a V. vulnificus promoter.